

SEQUENCE LISTING

<110> Soppet et al.

<120> G-Protein Parathyroid Hormone Receptor HLTG74

<130> PF201D1

<140> 09/236,468

<141> 1999-01-25

<150> 08/468,011

<151> 1995-06-06

<160> 28

<170> PatentIn Ver. 2.1

<210> 1

<211> 2003

<212> DNA

<213> Homo sapiens

<220>

<221> CDS

<222> (90)..(1715)

<400> 1

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Met Ala Trp Leu Gly Ala Ser Leu
1 5

cac gtc tgg ggt tgg cta atg ctc ggc agc tgc ctc ctg gcc aga gcc 161
His Val Trp Gly Trp Leu Met Leu Gly Ser Cys Leu Leu Ala Arg Ala
10 15 20

cag ctg gat tct gat ggc acc atc act ata gag gag cag att gtc ctt 209
Gln Leu Asp Ser Asp Gly Thr Ile Thr Ile Glu Glu Gln Ile Val Leu
25 30 35 40

gtg ctg aaa gcg aaa gta caa tgt gaa ctc aac atc aca gct caa ctc 257
Val Leu Lys Ala Lys Val Gln Cys Glu Leu Asn Ile Thr Ala Gln Leu
45 50 55

cag gag gga gaa ggt aat tgt ttc cct gaa tgg gat gga ctc att tgt 305
Gln Glu Gly Glu Asn Cys Phe Pro Glu Trp Asp Gly Leu Ile Cys
60 65 70

tgg ccc aga gga aca gtg ggg aaa ata tcg gct gtt cca tgc cct cct 353
Trp Pro Arg Gly Thr Val Gly Lys Ile Ser Ala Val Pro Cys Pro Pro
75 80 85

tat att tat gac ttc aac cat aaa gga gtt gct ttc cga cac tgt aac 401
Tyr Ile Tyr Asp Phe Asn His Lys Gly Val Ala Phe Arg His Cys Asn
90 95 100

ccc aat gga aca tgg gat ttt atg cac agc tta aat aaa aca tgg gcc 449
Pro Asn Gly Thr Trp Asp Phe Met His Ser Leu Asn Lys Thr Trp Ala
105 110 115 120

aat tat tca gac tgc ctt cgc ttt ctg cag cca gat atc agc ata gga	497		
Asn Tyr Ser Asp Cys Leu Arg Phe Leu Gln Pro Asp Ile Ser Ile Gly			
125	130	135	
aag caa gaa ttc tgt gaa cgc ctc tat gta atg tat acc gtt ggc tac	545		
Lys Gln Glu Phe Cys Glu Arg Leu Tyr Val Met Tyr Thr Val Gly Tyr			
140	145	150	
tcc atc tct ttt ggt tcc ttg gct gtg gct att ctc atc att ggt tac	593		
Ser Ile Ser Phe Gly Ser Leu Ala Val Ala Ile Leu Ile Ile Gly Tyr			
155	160	165	
ttc aga cga ttg cat tgc act agg aac tat atc cac atg cac tta ttt	641		
Phe Arg Arg Leu His Cys Thr Arg Asn Tyr Ile His Met His Leu Phe			
170	175	180	
gtg tct ttc atg ctg aga gct aca agc atc ttt gtc aaa gac aga gta	689		
Val Ser Phe Met Leu Arg Ala Thr Ser Ile Phe Val Lys Asp Arg Val			
185	190	195	200
gtc cat gct cac ata gga gta aag gag ctg gag tcc cta ata atg cag	737		
Val His Ala His Ile Gly Val Lys Glu Leu Glu Ser Leu Ile Met Gln			
205	210	215	
gat gac cca caa aat tcc att gag gca act tct gtg gac aaa tca caa	785		
Asp Asp Pro Gln Asn Ser Ile Glu Ala Thr Ser Val Asp Lys Ser Gln			
220	225	230	
tat atc ggg tgc aag att gct gtt gtg atg ttt att tac ttc ctg gct	833		
Tyr Ile Gly Cys Ile Ala Val Val Met Phe Ile Tyr Phe Leu Ala			
235	240	245	
aca aat tat tat tgg atc ctg gtg gaa ggt ctc tac ctg cat aat ctc	881		
Thr Asn Tyr Tyr Trp Ile Leu Val Glu Gly Leu Tyr Leu His Asn Leu			
250	255	260	
atc ttt gtg gct ttc ttt tcg gac acc aaa tac ctg tgg ggc ttc atc	929		
Ile Phe Val Ala Phe Phe Ser Asp Thr Lys Tyr Leu Trp Gly Phe Ile			
265	270	275	280
ttg ata ggc tgg ggg ttt cca gca gca ttt gtt gca gca tgg gct gtg	977		
Leu Ile Gly Trp Gly Phe Pro Ala Ala Phe Val Ala Ala Trp Ala Val			
285	290	295	
gca cga gca act ctg gct gat gcg agg tgc tgg gaa ctt agt gct gga	1025		
Ala Arg Ala Thr Leu Ala Asp Ala Arg Cys Trp Glu Leu Ser Ala Gly			
300	305	310	
gac atc aag tgg att tat caa gca ccg atc tta gca gct att ggg ctg	1073		
Asp Ile Lys Trp Ile Tyr Gln Ala Pro Ile Leu Ala Ala Ile Gly Leu			
315	320	325	
aat ttt att ctg ttt ctg aat acg gtt aga gtt cta gct acc aaa atc	1121		
Asn Phe Ile Leu Phe Leu Asn Thr Val Arg Val Leu Ala Thr Lys Ile			
330	335	340	
tgg gag acc aat gca gtt ggg cat gac aca agg aag caa tac agg aaa	1169		
Trp Glu Thr Asn Ala Val Gly His Asp Thr Arg Lys Gln Tyr Arg Lys			
345	350	355	360

ctg gcc aaa tcg aca ctg gtc ctg gtc cta gtc ttt gga gtg cat tac Leu Ala Lys Ser Thr Leu Val Leu Val Leu Val Phe Gly Val His Tyr 365 370 375	1217
atc gtg ttc gtg tgc ctg cct cac tcc ttc act ggg ctc ggg tgg gag Ile Val Phe Val Cys Leu Pro His Ser Phe Thr Gly Leu Gly Trp Glu 380 385 390	1265
atc cgc atg cac tgt gag ctc ttc aac tcc ttt cag ggt ttc ttt Ile Arg Met His Cys Glu Leu Phe Phe Asn Ser Phe Gln Gly Phe Phe 395 400 405	1313
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aag aag atg tgg agt cgg tgg aat ctc tcc gtg gac tgg aaa agg aca Lys Lys Met Trp Ser Arg Trp Asn Leu Ser Val Asp Trp Lys Arg Thr 425 430 435 440	1409
ccg cca tgt ggc agc cgc aga tgc ggc tca gtg ctc acc acc gtg acg Pro Pro Cys Gly Ser Arg Arg Cys Gly Ser Val Leu Thr Thr Val Thr 445 450 455	1457
cac agc acc agc cag tca cag gtg gcg gca gca cac gca tgg tgc His Ser Thr Ser Ser Gln Ser Gln Val Ala Ala Ala His Ala Trp Cys 460 465 470	1505
tta tct ctg gca aag ctg cca aga tcg cca gca gac agc ctg aca gcc Leu Ser Leu Ala Lys Leu Pro Arg Ser Pro Ala Asp Ser Leu Thr Ala 475 480 485	1553
aca tca ctt tac ctg gct atg tct gga gta act cag agc agg act gcc Thr Ser Leu Tyr Leu Ala Met Ser Gly Val Thr Gln Ser Arg Thr Ala 490 495 500	1601
tca cac act ctc tcc acg agg agc aac aag gaa gat agt ggg agg cag Ser His Thr Leu Ser Thr Arg Ser Asn Lys Glu Asp Ser Gly Arg Gln 505 510 515 520	1649
aga gat gat att cta atg gag aag cct tcc agg cct atg gaa tct aac Arg Asp Asp Ile Leu Met Glu Lys Pro Ser Arg Pro Met Glu Ser Asn 525 530 535	1697
cca gac act gaa gga tgacaaggag aaactgagga tggatctatgt atggacatgt Pro Asp Thr Glu Gly 540	1752
gtggctgact ttcatggc ggtccatgg ctgggtgtgt gagagggctt ggctgatact 1812	
cctatgctt agcacaaagg ctgaaaattc agttaaggtt ttacttaata atagttttta 1872	
ggctccatga attggctcct gtaaatacta acgacatgaa aatgcaagtg tcaatggagt 1932	
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<210> 2
<211> 541
<212> PRT
<213> Homo sapiens

<400> 2
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1 5 10 15
Gly Ser Cys Leu Leu Ala Arg Ala Gln Leu Asp Ser Asp Gly Thr Ile
20 25 30
Thr Ile Glu Glu Gln Ile Val Leu Val Leu Lys Ala Lys Val Gln Cys
35 40 45
Glu Leu Asn Ile Thr Ala Gln Leu Gln Glu Gly Glu Gly Asn Cys Phe
50 55 60
Pro Glu Trp Asp Gly Leu Ile Cys Trp Pro Arg Gly Thr Val Gly Lys
65 70 75 80
Ile Ser Ala Val Pro Cys Pro Pro Tyr Ile Tyr Asp Phe Asn His Lys
85 90 95
Gly Val Ala Phe Arg His Cys Asn Pro Asn Gly Thr Trp Asp Phe Met
100 105 110
His Ser Leu Asn Lys Thr Trp Ala Asn Tyr Ser Asp Cys Leu Arg Phe
115 120 125
Leu Gln Pro Asp Ile Ser Ile Gly Lys Gln Glu Phe Cys Glu Arg Leu
130 135 140
Tyr Val Met Tyr Thr Val Gly Tyr Ser Ile Ser Phe Gly Ser Leu Ala
145 150 155 160
Val Ala Ile Leu Ile Ile Gly Tyr Phe Arg Arg Leu His Cys Thr Arg
165 170 175
Asn Tyr Ile His Met His Leu Phe Val Ser Phe Met Leu Arg Ala Thr
180 185 190
Ser Ile Phe Val Lys Asp Arg Val Val His Ala His Ile Gly Val Lys
195 200 205
Glu Leu Glu Ser Leu Ile Met Gln Asp Asp Pro Gln Asn Ser Ile Glu
210 215 220
Ala Thr Ser Val Asp Lys Ser Gln Tyr Ile Gly Cys Lys Ile Ala Val
225 230 235 240
Val Met Phe Ile Tyr Phe Leu Ala Thr Asn Tyr Tyr Trp Ile Leu Val
245 250 255
Glu Gly Leu Tyr Leu His Asn Leu Ile Phe Val Ala Phe Phe Ser Asp
260 265 270
Thr Lys Tyr Leu Trp Gly Phe Ile Leu Ile Gly Trp Gly Phe Pro Ala
275 280 285
Ala Phe Val Ala Ala Trp Ala Val Ala Arg Ala Thr Leu Ala Asp Ala

290	295	300
Arg Cys Trp Glu Leu Ser Ala Gly Asp Ile Lys Trp Ile Tyr Gln Ala		
305	310	315
Pro Ile Leu Ala Ala Ile Gly Leu Asn Phe Ile Leu Phe Leu Asn Thr		
325	330	335
Val Arg Val Leu Ala Thr Lys Ile Trp Glu Thr Asn Ala Val Gly His		
340	345	350
Asp Thr Arg Lys Gln Tyr Arg Lys Leu Ala Lys Ser Thr Leu Val Leu		
355	360	365
Val Leu Val Phe Gly Val His Tyr Ile Val Phe Val Cys Leu Pro His		
370	375	380
Ser Phe Thr Gly Leu Gly Trp Glu Ile Arg Met His Cys Glu Leu Phe		
385	390	395
Phe Asn Ser Phe Gln Gly Phe Phe Val Ser Ile Ile Tyr Cys Tyr Cys		
405	410	415
Asn Gly Glu Val Gln Ala Glu Val Lys Lys Met Trp Ser Arg Trp Asn		
420	425	430
Leu Ser Val Asp Trp Lys Arg Thr Pro Pro Cys Gly Ser Arg Arg Cys		
435	440	445
Gly Ser Val Leu Thr Thr Val Thr His Ser Thr Ser Ser Gln Ser Gln		
450	455	460
Val Ala Ala Ala His Ala Trp Cys Leu Ser Leu Ala Lys Leu Pro Arg		
465	470	475
Ser Pro Ala Asp Ser Leu Thr Ala Thr Ser Leu Tyr Leu Ala Met Ser		
485	490	495
Gly Val Thr Gln Ser Arg Thr Ala Ser His Thr Leu Ser Thr Arg Ser		
500	505	510
Asn Lys Glu Asp Ser Gly Arg Gln Arg Asp Asp Ile Leu Met Glu Lys		
515	520	525
Pro Ser Arg Pro Met Glu Ser Asn Pro Asp Thr Glu Gly		
530	535	540

<210> 3

<211> 23

<212> DNA

<213> Artificial Sequence

<220>

<221> Primer_Bind

<223> This 5' primer sequence contains a SmaI restriction enzyme site followed by nucleotides corresponding to PTH receptor coding sequence.

<400> 3
cagccgtccc gggcttggcc tgg

23

<210> 4
<211> 27
<212> DNA
<213> Artificial Sequence

<220>
<221> Primer_Bind
<223> This 3' primer sequence contains a SalI restriction enzyme site and a sequence complementary to the human PTH receptor.

<400> 4
cctcagtgtc gacttgtcat ctttcag

27

<210> 5
<211> 27
<212> DNA
<213> Artificial Sequence

<220>
<221> Primer_Bind
<223> This 5' primer contains a HindIII restriction enzyme site and a nucleotide sequence corresponding to the 5' UTR of the cDNA encoding human PTH receptor.

<400> 5
gttggcatat tggaaagcttt ttgcggg

27

<210> 6
<211> 28
<212> DNA
<213> Artificial Sequence

<220>
<221> Primer_Bind
<223> This 3' primer sequence contains an XbaI restriction enzyme site, a translation stop codon, and nucleotides complementary to the human PTH receptor coding sequence.

<400> 6
cagtttcttag atgtcatcct tcagtgtc

28

<210> 7
<211> 39
<212> DNA
<213> Artificial Sequence

<220>
<221> Primer_Bind
<223> This 5' primer contains a SmaI restriction enzyme site, a nucleotide sequence to provide efficient initiation of translation in eukaryotic cells, and a nucleotide sequence corresponding to the human PTH receptor cDNA, including an initiation codon.

<400> 7
tcctaccggg gccggccatca tggcctggct gggggggcct

39

<210> 8
<211> 28
<212> DNA
<213> Artificial Sequence

<220>
<221> Primer_Bind
<223> This 3' primer contains an XbaI restriction enzyme site and a nucleotide sequence complementary to the 3' untranslated region of the PTH receptor cDNA.

<400> 8
cagttctag atgtcatcct tcagtgtc

28

<210> 9
<211> 60
<212> PRT
<213> Homo sapiens

<400> 9
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1 5 10 15

Lys Ser Gln Tyr Ile Gly Cys Lys Ile Ala Val Val Met Phe Ile Tyr
20 25 30

Phe Leu Ala Thr Asn Tyr Tyr Trp Ile Leu Val Glu Gly Leu Tyr Leu
35 40 45

His Asn Leu Ile Phe Val Ala Phe Phe Ser Asp Thr
50 55 60

<210> 10
<211> 60
<212> PRT
<213> Didelphis virginiana

<400> 10
Ile Thr Glu Glu Glu Leu Arg Ala Phe Thr Glu Pro Pro Pro Ala Asp
1 5 10 15

Lys Ala Gly Phe Val Gly Cys Arg Val Ala Val Thr Val Phe Leu Tyr
20 25 30

Phe Leu Thr Thr Asn Tyr Tyr Trp Ile Leu Val Glu Gly Leu Tyr Leu
35 40 45

His Ser Leu Ile Phe Met Ala Phe Phe Ser Glu Lys
50 55 60

<210> 11
<211> 60
<212> PRT
<213> Homo sapiens

<400> 11
Lys Tyr Leu Trp Gly Phe Ile Leu Ile Gly Trp Gly Phe Pro Ala Ala
1 5 10 15
Phe Val Ala Ala Trp Ala Val Ala Arg Ala Thr Leu Ala Asp Ala Arg
20 25 30
Cys Trp Glu Leu Ser Ala Gly Asp Ile Lys Trp Ile Tyr Gln Ala Pro
35 40 45
Ile Leu Ala Ala Ile Gly Leu Asn Phe Ile Leu Phe
50 55 60

<210> 12
<211> 60
<212> PRT
<213> *Didelphis virginiana*

<400> 12
Lys Tyr Leu Trp Gly Phe Thr Leu Phe Gly Trp Gly Leu Pro Ala Val
1 5 10 15
Phe Val Ala Val Trp Val Thr Val Arg Ala Thr Leu Ala Asn Thr Glu
20 25 30
Cys Trp Asp Leu Ser Ser Gly Asn Lys Lys Trp Ile Ile Gln Val Pro
35 40 45
Ile Leu Ala Ala Ile Val Val Asn Phe Ile Leu Phe
50 55 60

<210> 13
<211> 52
<212> PRT
<213> Homo sapiens

<400> 13
Leu Asn Thr Val Arg Val Leu Ala Thr Lys Ile Trp Glu Thr Asn Ala
1 5 10 15
Val Gly His Asp Thr Arg Lys Gln Tyr Arg Lys Leu Ala Lys Ser Thr
20 25 30
Leu Val Leu Val Leu Val Phe Gly Val His Tyr Ile Val Phe Val Cys
35 40 45
Leu Pro His Ser
50

<210> 14
<211> 52
<212> PRT
<213> *Didelphis virginiana*

<400> 14
Ile Asn Ile Ile Arg Val Leu Ala Thr Lys Leu Arg Glu Thr Asn Ala
1 5 10 15

Gly Arg Cys Asp Thr Arg Gln Gln Tyr Arg Lys Leu Leu Lys Ser Thr
20 25 30

Leu Val Leu Met Pro Leu Phe Gly Val His Tyr Ile Val Phe Met Ala
35 40 45

Thr Pro Tyr Thr
50

<210> 15
<211> 60
<212> PRT
<213> *Homo sapiens*

<400> 15
Glu Gly Asn Cys Phe Pro Glu Trp Asp Gly Leu Ile Cys Trp Pro Arg
1 5 10 15

Gly Thr Val Gly Lys Ile Ser Ala Val Pro Cys Pro Pro Tyr Ile Tyr
20 25 30

Asp Phe Asn His Lys Gly Val Ala Phe Arg His Cys Asn Pro Asn Gly
35 40 45

Thr Trp Asp Phe Met His Ser Leu Asn Lys Thr Trp
50 55 60

<210> 16
<211> 60
<212> PRT
<213> *Didelphis virginiana*

<400> 16
Asp Gly Phe Cys Leu Pro Glu Trp Asp Asn Ile Val Cys Trp Pro Ala
1 5 10 15

Gly Val Pro Gly Lys Val Val Ala Val Pro Cys Pro Asp Tyr Ile Tyr
20 25 30

Asp Phe Asn His Lys Gly Arg Ala Tyr Arg Arg Cys Asp Ser Asn Gly
35 40 45

Ser Trp Glu Leu Val Pro Gly Asn Asn Arg Thr Trp
50 55 60

<210> 17
<211> 10
<212> PRT
<213> Homo sapiens

<400> 17
Ala Asn Tyr Ser Asp Cys Leu Arg Phe Leu
1 5 10

<210> 18
<211> 10
<212> PRT
<213> Didelphis virginiana

<400> 18
Ala Asn Tyr Ser Glu Cys Val Lys Phe Leu
1 5 10

<210> 19
<211> 60
<212> PRT
<213> Homo sapiens

<400> 19
Lys Gln Glu Phe Cys Glu Arg Leu Tyr Val Met Tyr Thr Val Gly Tyr
1 5 10 15

Ser Ile Ser Phe Gly Ser Leu Ala Val Ala Ile Leu Ile Ile Gly Tyr
20 25 30

Phe Arg Arg Leu His Cys Thr Arg Asn Tyr Ile His Met His Leu Phe
35 40 45

Val Ser Phe Met Leu Arg Ala Thr Ser Ile Phe Val
50 55 60

<210> 20
<211> 60
<212> PRT
<213> Didelphis virginiana

<400> 20
Glu Arg Glu Val Phe Asp Arg Leu Gly Met Ile Tyr Thr Val Gly Tyr
1 5 10 15

Ser Ile Ser Leu Gly Ser Leu Thr Val Ala Val Leu Ile Leu Gly Tyr
20 25 30

Phe Arg Arg Leu His Cys Thr Arg Asn Tyr Ile His Met His Leu Phe
35 40 45

Val Ser Phe Met Leu Arg Ala Val Ser Ile Phe Ile
50 55 60

<210> 21
<211> 21
<212> PRT
<213> Homo sapiens

<400> 21
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1 5 10 15
Leu Ile Met Gln Asp
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<210> 22
<211> 21
<212> PRT
<213> *Didelphis virginiana*

<400> 22
Lys Asp Ala Val Leu Tyr Ser Gly Val Ser Thr Asp Glu Ile Glu Arg
1 5 10 15
Ile Thr Glu Glu Glu
20

<210> 23
<211> 59
<212> PRT
<213> Homo sapiens

<400> 23
Thr Gly Leu Gly Trp Glu Ile Arg Met His Cys Glu Leu Phe Phe Asn
1 5 10 15
Ser Phe Gln Gly Phe Phe Val Ser Ile Ile Tyr Cys Tyr Cys Asn Gly
20 25 30
Glu Val Gln Ala Glu Val Lys Lys Met Trp Ser Arg Trp Asn Leu Ser
35 40 45
Val Asp Trp Lys Arg Thr Pro Pro Cys Gly Ser
50 55

<210> 24
<211> 59
<212> PRT
<213> *Didelphis virginiana*

<400> 24
Ser Gly Ile Leu Trp Gln Val Gln Met His Tyr Glu Met Leu Phe Asn
1 5 10 15
Ser Phe Gln Gly Phe Phe Val Ala Ile Ile Tyr Cys Phe Cys Asn Gly
20 25 30
Glu Val Gln Ala Glu Ile Lys Lys Ser Trp Ser Arg Trp Thr Leu Ala
35 40 45

Leu Asp Phe Lys Arg Lys Ala Arg Ser Gly Ser
50 55

<210> 25
<211> 37
<212> PRT
<213> Homo sapiens

<400> 25
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1 5 10 15

Leu Val Leu Lys Ala Lys Val Gln Cys Glu Leu Asn Ile Thr Ala Gln
20 25 30

Leu Gln Glu Gly Glu
35

<210> 26
<211> 37
<212> PRT
<213> Didelphis virginiana

<400> 26
Ala Leu Val Asp Ala Asp Asp Val Ile Thr Lys Glu Glu Gln Ile Ile
1 5 10 15

Leu Leu Arg Asn Ala Gln Ala Gln Cys Glu Gln Arg Leu Lys Glu Val
20 25 30

Leu Arg Val Pro Glu
35

<210> 27
<211> 23
<212> PRT
<213> Homo sapiens

<400> 27
Ile Ser Gly Lys Ala Ala Lys Ile Ala Ser Arg Gln Pro Asp Ser His
1 5 10 15

Ile Thr Leu Pro Gly Tyr Val
20

<210> 28
<211> 23
<212> PRT
<213> Didelphis virginiana

<400> 28
Leu Ser Pro Arg Leu Ala Pro Gly Ala Gly Ala Ser Ala Asn Gly His
1 5 10 15

His Gln Leu Pro Gly Tyr Val
20